

- a) BLC = RP-II protease from *Bacillus licheniformis*,  
 b) AA513 = RP-II protease from *Bacillus halmapalus* AA513  
 c) AC116 = RP-II protease from *Bacillus licheniformis* AC116  
 d) BO32 = RP-II protease from *Bacillus pumilus* BO32  
 e) CDJ31 = RP-II protease from *Bacillus licheniformis* CDJ31  
 f) JA96 = RP-II protease from *Bacillus pumilus* JA96  
 g) MPR = RP-II protease from *Bacillus subtilis* IS75

- a) 1 SVIGSDDRTRVTNTTAYPYRAIVHIS...SSIGSCTGWMIGPKTVATAG 46  
 b) VVIGDDGRRQVQNTSFMPFRALTYIEFGNLTSTWSCSGGVIGTDLVVTNA
- a) 47 HCIYDTSSGSFAGTATVSPGRNGTSYPYGSVKSTRYFIPSGWR.SGNTNY 95  
 b) HCV...EGSVL.AGTVVPGMNSQWAYGHYRVTQIIYPDQYRNGASEF
- a) 96 DYGAIELS...EPIGNTVGYFGYSYTTSSLVGTTVTISGYPGDKTAGT 140  
 b) DYAILRVAPDSDGRHIGNRAGILSFTETGTVNENTFLRTYGYPGDKISET
- a) 141 Q...WQHSG.PIATSEYTKLQYAMDYGGQSGSPVFEQSSSRTNCSGPC 185  
 b) KLISLWGMVGRSDAFLHRDLLFYNMPTYFGQSGSPVLNSVDSMVAVHNAG
- a) 186 SLAVHTNGVYGGSSYNRGTRITKEVFDNLTNWKNNSAQ 222  
 b) YIVGGNREINGGPKIRRDFTNLFNQMN.....
- a) 1 SVIGSDDRTRVTNTTAYPYRAIVHISSSIGSCTGWMIGPKTVATAGHCIY 50  
 c) SVIGSDERTRVTDTTAFPYRAIVHISSSIGSCTGWLIGPKTVATAGHCVY
- a) 51 DTSSGSFAGTATVSPGRNGTSYPYGSVKSTRYFIPSGWRSGNTNYDYGAI 100  
 c) DTASRSFAGTATVSPGRNGSAYPYGSVTSTRYFIPSGWQSGNSNYDYAAI
- a) 101 ELSEPIGNTVGYFGYSYTTSSLVGTTVTISGYPGDKTAGTQWQHSGPIAI 150  
 c) ELSQPIGNTVGYFGYSYTASSLAGAGVTISGYPGDKTTGTQWQMSGTIIV
- a) 151 SETYKLQYAMDYGGQSGSPVFEQSSSRTNCSGPCSLAVHTNGVYGGSSY 200  
 c) SETYKLQYAITDYGGQSGSPVYEKSSSRTNCSGPCSLAVHTNGVYGGSSY
- a) 201 NRGTRITKEVFDNLTNWKNNSAQ 222  
 c) NRGTRITKEVFDNFTSWKNNSAQ 222

Fig. 1a

a) 1 SVIGSDDRTRVTNTTAYPYRAIVHISSSIGSCTGWMIGPKTVATAGHCIY 50  
d) VVIGDDGRTKVANTRVAPYNSIAYTTFGGSSCTGTLIAPNKILTNGHCVY

a) 51 DTSSGSFAGTATVSPGRNGTSYPYGSVKSTRYFIPSGW.RSGNTNYDYGA 99  
d) NTASRSYSAGSVYPMNDSTAVNGSANMTEFYVPSGYINTGASQYDFAV

a) 100 IELSEPIGNTVGYFGYSYTTSSLVGTTVTISGYPGDKTAGT....QWQHS 145  
d) IKTDTNIGNTVGYRSIRQVT.NLTGTTIKISGYPGDKMRSTGKISQWEMS

a) 146 GPIAISETYKLQYAMDYGGQSGSPVFEQSSSRTNC.SGPCSLAVHTNGV 194  
d) GPVTREDTNLAYYMIDTFSGNSGSAMLDQNQQIVGVHNAGYSNGTINGGP

a) 195 YGGSSYNRGTRITKEVFDNLTNWKNFAQ 222  
d) KATAAFVEFINYAKAQ.....

a) 1 SVIGSDDRTRVTNTTAYPYRAIVHISSSIGSCTGWMIGPKTVATAGHCIY 50  
e) SVIGSDERTRVTNTTAYPYRAIVHISSSIGSCTGSLIGPKTVATAGHCIY

a) 51 DTSSGSFAGTATVSPGRNGTSYPYGSVKSTRYFIPSGWRSGNTNYDYGAI 100  
e) DTASGSFAGTATVSPGRNGSTYPYGSVTSTRYFIPSGYRSGNSNYDYGAI

a) 101 ELSEPIGNTVGYFGYSYTTSSLVGTTVTISGYPGDKTAGTQWQHSGPIAI 150  
e) ELSQPIGNTVGYFGYSYTTSSLVGSSVTIIGYPGDKTSQTQWQMSGNIIV

a) 151 SETYKLQYAMDYGGQSGSPVFEQSSSRTNC.SGPCSLAVHTNGVYGGSSY 200  
e) SETYKLQYAITDYGGQSGSPVYEASSSRTNC.SGPCSLAVHTNGVYGGSSY

a) 201 NRGTRITKEVFDNLTNWKNFAQ 222  
e) NRGTRITKEVFDNLTNWKNFAQ

a) 1 SVIGSDDRTRVTNTTAYPYRAIVHISSSIGSCTGWMIGPKTVATAGHCIY 50  
f) VVIGDDGRTKVTNTRVAPYNSIAYITFGGSSCTGTLIAPNKILTNGHCVY

a) 51 DTSSGSFAGTATVSPGRNGTSYPYGSVKSTRYFIPSGW.RSGNTNYDYGA 99  
f) NTATRSYSAGSVYPMNDSTAVNGSANMTEFYVPSGYINTGASQYDFAV

a) 100 IELSEPIGNTVGYFGYSYTTSSLVGTTVTISGYPGDKTAGT....QWQHS 145  
f) IKTDTNIGNTVGYRSIRQVT.NLTGTTIKISGYPGDKMRSTGKVSQWEMS

a) 146 GPIAISETYKLQYAMDYGGQSGSPVFEQSSSRTNC.SGPCSLAVHTNGV 194  
f) GPVTREDTNLAYYTIDTFSGNSGSAMLDQNQQIVGVHNAGYSNGTINGGP

a) 195 YGGSSYNRGTRITKEVFDNLTNWKNFAQ 222  
f) KATAAFVEFINYAKAQ.....

Fig. 1b

a) 1 SVIGSDDRTRVTNTTAYPYRAIVHIS.....SSIGSCTGWMIGPKTVAT 44  
g) SIIGTDERTRISSTTSFPYRATVQLSIKYPNTSSSTYGCTGFLVNPNTVVT

a) 45 AGHCIYDTSSGSFAGTATVSPGRNGTSYPYGSVKSTRYFIPSGW.RSGNT 93  
g) AGHCVYSQDHG.WASTITAAPGRNGSSYPYGTYSGTMFYSVKGWTESKDT

a) 94 NYDYGAIELSEPIGNTVGYFGYSYT.TSSLVGTTVTISGYPGDKTAGTQW 142  
g) NYDYGAIKLNGSPGNTVGWYGYRTTNSSSPVGLSSSVTGFPCKDTFGTMW

a) 143 QHSGPIAISETYKLQYAMDYGGQSGSPVFEQSSSRNTNCSGPCSLAVHTN 192  
g) SDTKPIRSAETYKLTYTTDTYGCQSGSPVYRNYSD....TGQTAIAIHTN

a) 193 GVGSSSYNRGTRITKEVFDNLTNWKNQAQ 222  
g) ...GGSSYNLGTTRVTNDVFNNIQYWANQ..

Fig. 1c

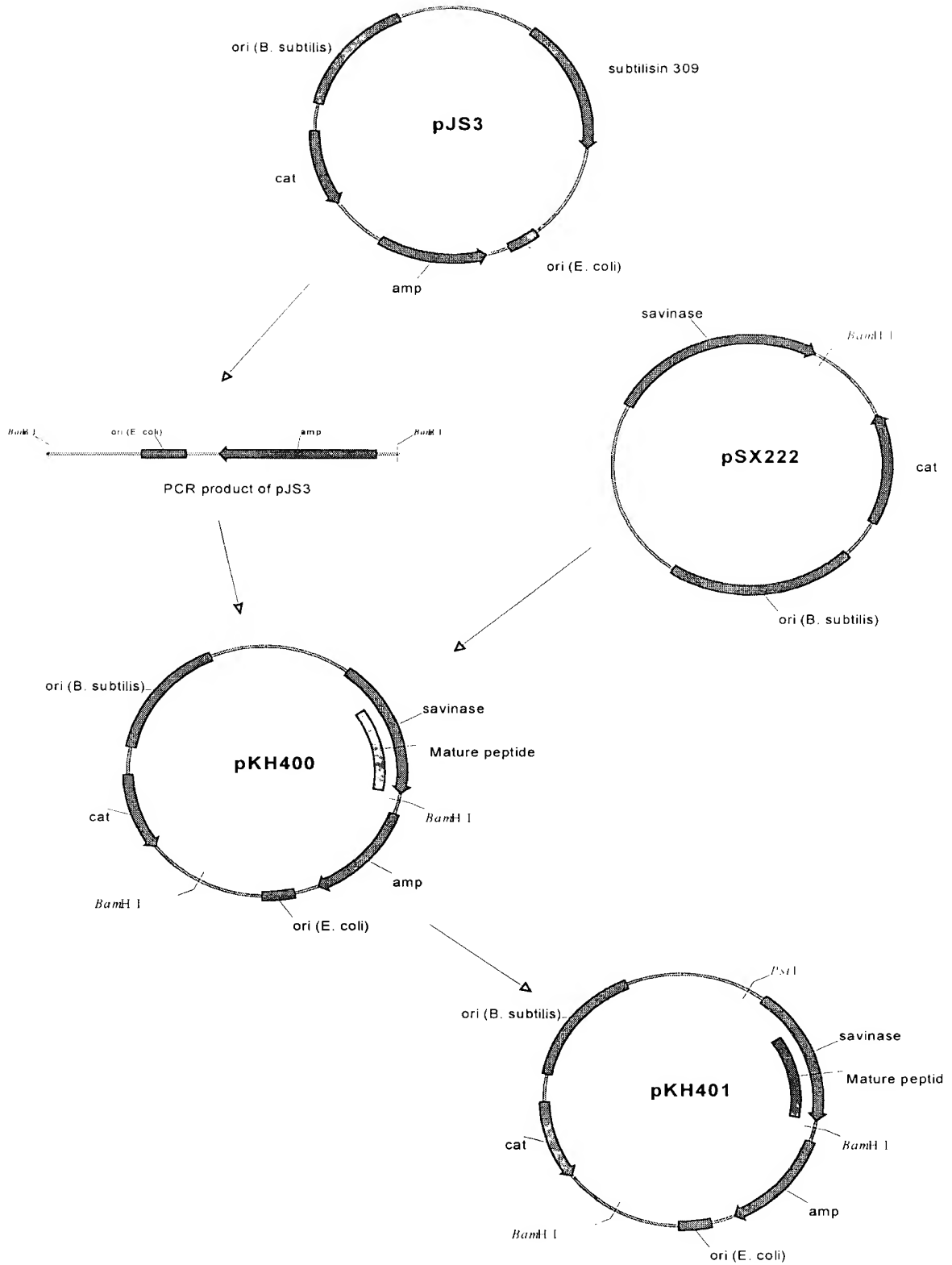


Fig 2a

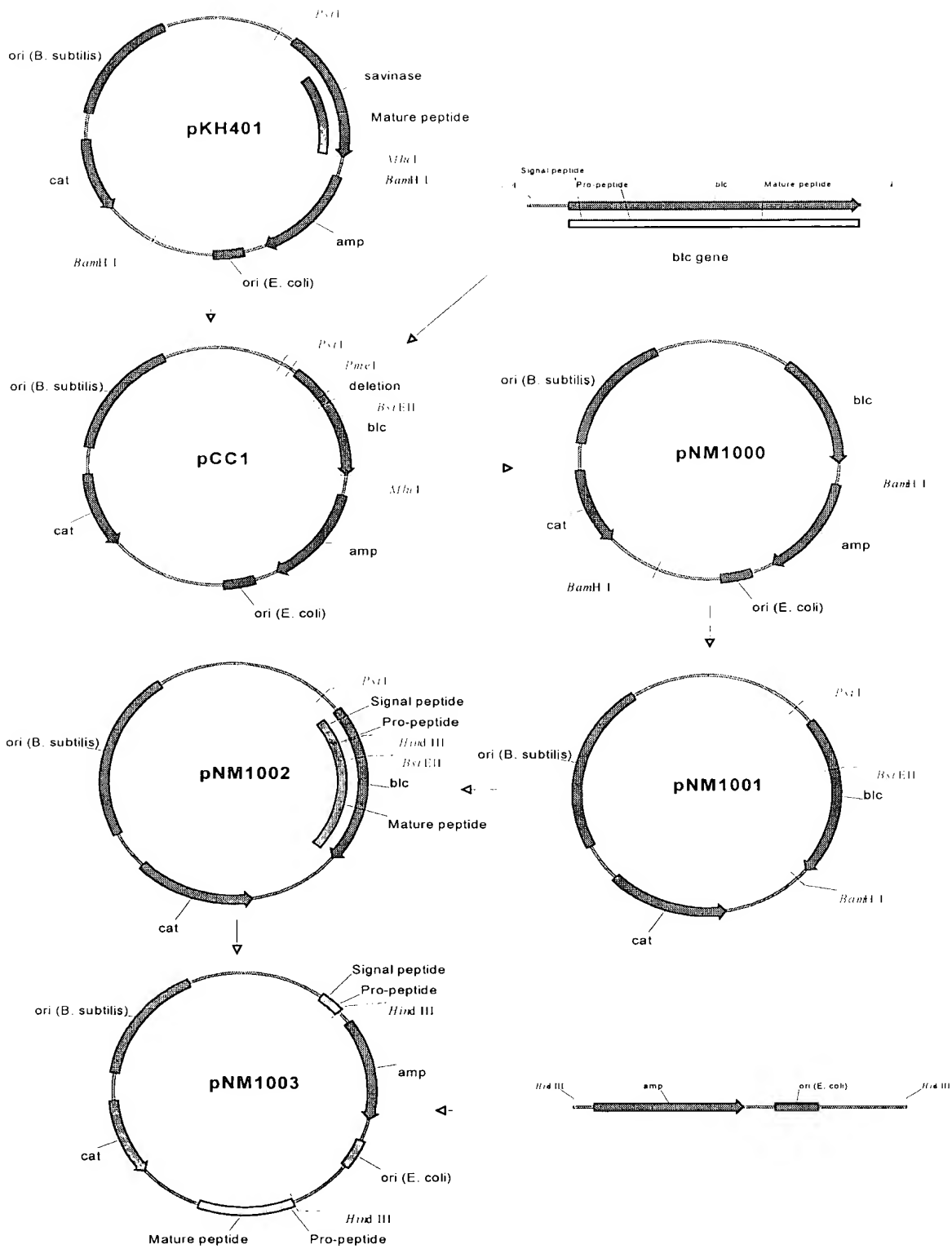


Fig 2b